

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/549,441
Source: IFWP
Date Processed by STIC: 8/18/06

ENTERED



IFWP

RAW SEQUENCE LISTING DATE: 08/18/2006
PATENT APPLICATION: US/10/549,441 TIME: 15:04:50

Input Set : A:\51471-20016.00.txt
Output Set: N:\CRF4\08182006\J549441.raw

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4 <110> APPLICANT: Shelton, David L.
6 <120> TITLE OF INVENTION: METHODS FOR TREATING TAXOL-INDUCED GUT
7 DISORDER
9 <130> FILE REFERENCE: 514712001600
11 <140> CURRENT APPLICATION NUMBER: 10/549,441
C--> 12 <141> CURRENT FILING DATE: 2005-09-16
14 <150> PRIOR APPLICATION NUMBER: PCT/US2004/008865
15 <151> PRIOR FILING DATE: 2004-03-22
17 <150> PRIOR APPLICATION NUMBER: US 60/456,648
18 <151> PRIOR FILING DATE: 2003-03-20
20 <160> NUMBER OF SEQ ID NOS: 10
22 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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25 <211> LENGTH: 123
26 <212> TYPE: PRT
27 <213> ORGANISM: Artificial Sequence
29 <220> FEATURE:
30 <223> OTHER INFORMATION: Synthetic Construct
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33 Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
34 1 5 10 15
35 Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Ser Tyr
36 20 25 30
37 Arg Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
38 35 40 45
39 Gly Glu Ile Tyr Pro Ser Asn Ala Arg Thr Asn Tyr Asn Glu Lys Phe
40 50 55 60
41 Lys Ser Arg Val Thr Met Thr Arg Asp Thr Ser Thr Ser Thr Val Tyr
42 65 70 75 80
43 Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
44 85 90 95
45 Ala Arg Lys Tyr Tyr Gly Asn Thr Arg Arg Ser Trp Tyr Phe Asp
46 100 105 110
47 Val Trp Gly Gln Gly Thr Thr Val Thr Val Ser
48 115 120
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52 <211> LENGTH: 113
53 <212> TYPE: PRT
54 <213> ORGANISM: Artificial Sequence
56 <220> FEATURE:
57 <223> OTHER INFORMATION: Synthetic Construct
59 <400> SEQUENCE: 2
60 Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly

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61 1 5 10 15
62 Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Glu Ser Ile Asp Asn Tyr
63 20 25 30
64 Gly Ile Ser Phe Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro
65 35 40 45
66 Lys Leu Leu Ile Tyr Ala Ala Ser Asn Arg Gly Ser Gly Val Pro Ser
67 50 55 60
68 Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Phe Thr Ile Ser
69 65 70 75 80
70 Ser Leu Gln Pro Glu Asp Ile Ala Thr Tyr Tyr Cys Gln Gln Ser Lys
71 85 90 95
72 Thr Val Pro Arg Thr Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys Arg
73 100 105 110
74 Thr
78 <210> SEQ ID NO: 3
79 <211> LENGTH: 13
80 <212> TYPE: PRT
81 <213> ORGANISM: Artificial Sequence
83 <220> FEATURE:
84 <223> OTHER INFORMATION: Synthetic Construct
86 <400> SEQUENCE: 3
87 Lys Ala Ser Gly Tyr Thr Phe Thr Ser Tyr Arg Ile His
88 1 5 10
91 <210> SEQ ID NO: 4
92 <211> LENGTH: 17
93 <212> TYPE: PRT
94 <213> ORGANISM: Artificial Sequence
96 <220> FEATURE:
97 <223> OTHER INFORMATION: Synthetic Construct
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100 Glu Ile Tyr Pro Ser Asn Ala Arg Thr Asn Tyr Asn Glu Lys Phe Lys
101 1 5 10 15
102 Ser
106 <210> SEQ ID NO: 5
107 <211> LENGTH: 18
108 <212> TYPE: PRT
109 <213> ORGANISM: Artificial Sequence
111 <220> FEATURE:
112 <223> OTHER INFORMATION: Synthetic Construct
114 <400> SEQUENCE: 5
115 Cys Ala Arg Lys Tyr Tyr Gly Asn Thr Arg Arg Ser Trp Tyr Phe
116 1 5 10 15
117 Asp Val
121 <210> SEQ ID NO: 6
122 <211> LENGTH: 15
123 <212> TYPE: PRT
124 <213> ORGANISM: Artificial Sequence
126 <220> FEATURE:
127 <223> OTHER INFORMATION: Synthetic Construct

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129 <400> SEQUENCE: 6
130 Arg Ala Ser Glu Ser Ile Asp Asn Tyr Gly Ile Ser Phe Leu Ala
131 1 5 10 15
134 <210> SEQ ID NO: 7
135 <211> LENGTH: 7
136 <212> TYPE: PRT
137 <213> ORGANISM: Artificial Sequence
139 <220> FEATURE:
140 <223> OTHER INFORMATION: Synthetic Construct
142 <400> SEQUENCE: 7
143 Ala Ala Ser Asn Arg Gly Ser
144 1 5
147 <210> SEQ ID NO: 8
148 <211> LENGTH: 9
149 <212> TYPE: PRT
150 <213> ORGANISM: Artificial Sequence
152 <220> FEATURE:
153 <223> OTHER INFORMATION: Synthetic Construct
155 <400> SEQUENCE: 8
156 Gln Gln Ser Lys Thr Val Pro Arg Thr
157 1 5
160 <210> SEQ ID NO: 9
161 <211> LENGTH: 654
162 <212> TYPE: DNA
163 <213> ORGANISM: Artificial Sequence
165 <220> FEATURE:
166 <223> OTHER INFORMATION: Synthetic Construct
168 <400> SEQUENCE: 9
169 gatatccaga tgacacagtc cccatcctcc ctgtctgcct ctgtgggtga ccgcgtcacc 60
170 atcacctgcc gcgcaagtga gagcatcgac aactatggca ttcccttccct ggcttgtat 120
171 cagcagaagc cgggcaaagc accaaaactc ctgatctatg ctgcatcaa tcggggttca 180
172 ggtgtcccat cacgcttcag tggcagtggc tctggtag agattcacctt caccattagc 240
173 agcctgcaac cagaagatata tggccacttat tactgccaac agagtaagac tgtgccacgc 300
174 actttcgggtc aaggcaccaa gctggagatc aaacgcactg tggctgcacc atctgtcttc 360
175 atcttccttc catctgatga gcagttaaaa tccggaaactg cctctgttgt gtgcctgctg 420
176 aataacttct atccacgcga gccaagatc cagtggaagg tggataacgc cctccaatcc 480
177 ggtaactccc aggagagtgt cacagagcag gacagcaagg acagcaccta cagcctcagc 540
178 agcaccctga ccctgagcaa agcagactac gagaacaca aagtctacgc ctgcgaagcc 600
179 acccatcagg gcctgagttc tccagtcaca aagagctca accgcggta gtgc 654
181 <210> SEQ ID NO: 10
182 <211> LENGTH: 1350
183 <212> TYPE: DNA
184 <213> ORGANISM: Artificial Sequence
186 <220> FEATURE:
187 <223> OTHER INFORMATION: Synthetic Construct
189 <400> SEQUENCE: 10
190 caggtgcagtc tggtgctgag gtgaagaagc ctggcgcttc cgtgaaggtt 60
191 tcctgcaaag catctggta caccttacc agctatcgga tccactgggt gcgccaagcc 120
192 cctggtaag gcctggagtg gatggcgaa atctacccaa gcaacgcgcg cactaactac 180

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193 aacgagaagt tcaaatacccg ggtgaccatg actcgcgata cctccaccag cactgtctac 240
194 atggaaactga gctctctgcg ctctgaggac actgctgtgt attactgtgc ccgcaagtac 300
195 tattacggca atacgcgtcg ctccctggta ttcgatgtgt ggggccagg taccactgtt 360
196 accgtgtccct ctgcctccac caagggccca tctgtcttcc cactggcccc atgctcccgc 420
197 agcacccctcg agagcacagc cgccctgggc tgccctggta aggactactt cccagaacct 480
198 gtgaccgtgt cctggaaactc tggcgctctg accagcggcg tgcacacett cccagctgtc 540
199 ctgcagtccct caggctctcta ctccctcagc agcgtggta ccgtgccatc cagcaacttc 600
200 ggcacccaga cctacacactg caacgttagat cacaagccaa gcaacaccaa ggtcgacaag 660
201 accgtggaga gaaagtgttg tgtggagtgt ccaccttgta cagcccttcc agtggccgga 720
202 ccatccgtgt tcctgttccc tccaaagcca aaggacaccc ttagtatctc cagaacccca 780
203 gaggtgaccc ttgtgtgtgt ggacgtgtcc cacgaggacc cagaggtgca gttcaactgg 840
204 tatgtggacg gagtggaggt gcacaacgcc aagaccaagc caagagagga gcagttcaac 900
205 tccacccatca gagtgggtgag cgtgctgacc gtgggtgcacc aggactggct gaacggaaag 960
206 gaggataagt gtaagggtgtc caacaaggga ctggccatcca gcatcgagaa gaccatctcc 1020
207 aagaccaagg gacagccaaag agagccacag gtgtataccct tgccaccatc cagagaggag 1080
208 atgaccaaga accaggtgtc cctgacctgt ctgggtgaagg gattctatcc atccgacatc 1140
209 gccgtggagt gggagtc当地 cggacagccca gagaacaact ataagaccac ccctccaatg 1200
210 ctggactccg acggatcctt tttccctgtat tccaaagctga ccgtggacaa gtccagatgg 1260
211 cagcaggaa acgtgttctc ttgttccgtg atgcacgagg ccctgcacaa ccactataacc 1320
212 cagaagagcc tggccctgtc tccaggaaag 1350

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/549,441

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Input Set : A:\51471-20016.00.txt

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L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date